

Amendments to the Specification:

Please amend Table 3 on page 33 as follows:

Table 3. Inferred Amino Acid Sequence of the R2 Envelope Clone from Donor 2

Amino Acid Residue ²					Residue
MRVKGIRRNY	QHWGWGTML	LGLLMICSAT	EKLWVTVYYG	VPVWKEATTT	50
LFCASDAKAY	DTEAHNVWAT	HACVPTDPNP	QEVELNVNTE	NFNMWKNNMV	100
EQMHEDIISL	WDQSLKPCVK	LTPLCVTLNC	TDLRNTTNTN	NSTDNNNSNS	150
EGTIKGEMK	NCSFNAITSI	GDKMQKEYAL	LYKLDIEPID	NDNTSYRLIS	200
CNTSVITQAC	PKISFEPIPI	HYCAPAGFAI	LKCNDKKFSG	KGSCKNVSTV	250
QCTHGIRPVV	STQLLLNGSL	AEEEVVIRSE	NFTNNAKTII	VQLREPVKIN	300
CSRPNNNTRK	SIPMGPGRAF	YTTGQIIGDI	RQAHCNISKT	NWTNALKQVV	350
EKLGEQFNKT	KIVFTNSSGG	DPEIVTHSFN	CAGEFFYCNT	TQLFDSIWNS	400
ENGTWNITRG	LNNTGRNDTI	TLPCRKQII	NRWQEVGKAM	YAPPIKGNIS	450
CSSNITGLLL	TRDGGKDDNS	RDGNETFRPG	GGDMRDNWS	ELYKYKVVKI	500
EPLGVAPTKA	KRRVVQREER	AVGLGAMFIG	FLGAAGSTMG	AASVTLTVQA	550
RQLLSGIVQQ	QSNLLRAIEA	QQHLLQLTVW	GIKQLQARIL	AVERYLKDQQ	600
LLGIWGCSGK	LICTTTVPWN	ASWSKNKTLE	AIWNNMTWMQ	WDKEIDNYTK	650
				WDKEIDNYTS	
LIYSLIEESQ	IQQEKNEQEL	LELDKWANKW	NWFDISNWLW	YIKIFIMIVG	700
GLVGLRIVFV	VLSIVNRVRQ	GYSPLSFQTR	LPAPRGPDPR	EEIEEEGGDR	750
DRDRSGLLVD	GFLTLIWVDL	RSLCLFSYHR	LRDLLLIVTR	IVELLGRRGW	800
EILKYWWNLL	QYWSQELKNS	AVSLFNATAI	AVAEGTDRVI	EVLQRVGRAL	850
LHIPTRIRQG	LERALL				866

²Amino Acid Residues are identified by standard single letter designations. Predicted N-linked glycosylation sites are indicated by shading and bolding